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DATE MAILED: 12/11/2006

APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.	
10/088,724	06/14/2002	Ikuo Nishimoto	082377-00000US	6929	
7590 12/11/2006		006	EXAM	EXAMINER	
Joe Liebeschuetz			CHERNYSHEV, OLGA N		
Townsend & 7	ownsend & Crew				
8th Floor			ART UNIT	PAPER NUMBER	
Two Embarcadero Center			1649		
Can Consider	CA 04111 2024				

Please find below and/or attached an Office communication concerning this application or proceeding.

U.S. Patent and Trademark Office PTOL-326 (Rev. 08-06)

Paper No(s)/Mail Date _

Information Disclosure Statement(s) (PTO/SB/08)

5) Notice of Informal Patent Application

6) Other: copy of the sequence alignment.

DETAILED ACTION

Response to amendment

1. Applicant's submission after final filed on October 31, 2006 has been entered. Claim 5 has been amended as requested in the amendment filed on October 31, 2006. Following the amendment, claims 1, 2, 4-8, 13, 20-22, 27-30, 35-38, 43 and 45 are pending in the instant application.

Claims 1, 2, 4-8, 13, 20-22, 27-30, 35-38, 43 and 45 are under examination in the instant office action.

2. During further examination and consideration, the relevant art of record has been discovered. Therefore, the finality of the previous Office action is withdrawn and the new grounds of rejection(s) are as follows.

Claim Rejections - 35 USC § 102

3. The following is a quotation of the appropriate paragraphs of 35 U.S.C. 102 that form the basis for the rejections under this section made in this Office action:

A person shall be entitled to a patent unless -

- (b) the invention was patented or described in a printed publication in this or a foreign country or in public use or on sale in this country, more than one year prior to the date of application for patent in the United States.
- 4. Claims 1, 5, 6, 7, 8, 13, 20, 28 and 36 are rejected under 35 U.S.C. 102(b) as being anticipated by Paznekas et al., 1997 (Biochem. Biophys. Res. Comm., 238, pp.1-6) and Dixon et al., 1997 (Hum. Mol. Gen., 6, No. 5, pp.727-37).

Claims 1, 5, 6, 7, 8, 13, 20, 28, 36 and 63 encompass a polypeptide of SEQ ID NO: 63, polynucleotide encoding a polypeptide of SEQ ID NO: 63, vectors, host cells and methods of

Art Unit: 1649

recombinant production of the polypeptide of SEQ ID NO: 63. Publications of Paznekas et al. and Dixon et al. disclose polypeptide and cDNA sequences, which have 100% identity to the instant claimed polypeptide of SEQ ID NO: 63 and encoding DNA, see copy of the sequence alignment attached to the instant office action. Thus the claimed subject matter of claims 1, 5, 6, 7, 8, 13, 20, 28, 36 and 63 is fully anticipated by these references.

Claim Objections

Claims 2, 4, 22, 27, 30, 35, 38, 43 and 45 are objected to for reciting non-elected subject 5. matter. Applicant's attention is directed to paper filed on October 16, 2003, in which election of a polypeptide of SEQ ID NO: 5 was made. During further examination of the instant application, polypeptide of SEQ ID NO: 5 was found to be novel, therefore the sequence search of the generic sequence of a polypeptide of SEQ ID NO: 63 was made and at that time the polypeptide of SEQ ID NO: 63 was considered free of prior art. Because claim 1 was considered free of prior art, dependent claims reciting plurality of sequences, were randomly searched and consequently considered free of prior art and allowable. However, at present, in view of the newly discovered references disclosing the polypeptide of the generic claim (SEQ ID NO: 63, see rejection of record in section 4 of the instant office action), only polypeptides of SEQ ID NO: 5, 33 and 101, as elected and previously searched, but not the full scope of the claims reciting plurality of different sequences, are considered to be allowable.

Art Unit: 1649

Conclusion

6. Claims 1, 5, 6, 7, 8, 13, 20, 28 and 36 are rejected. Claims 2, 4, 22, 27, 30, 35, 38, 43 and 45 are objected to. Claims 21, 29 and 37 are objected to for being dependent from rejected claims but would be allowed if rewritten in independent form.

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Olga N. Chernyshev whose telephone number is (571) 272-0870. The examiner can normally be reached on 8:00 AM to 5:00 PM.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Janet L. Andres can be reached on (571) 272-0867. The fax phone number for the organization where this application or proceeding is assigned is 571-273-8300.

Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see http://pair-direct.uspto.gov. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free). If you would like assistance from a USPTO Customer Service Representative or access to the automated information system, call 800-786-9199 (IN USA OR CANADA) or 571-272-1000.

Art Unit 1649

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RESULT 7
TCOF_MOUSE
     TCOF MOUSE
                      STANDARD;
                                       PRT; 1320 AA.
AC
     008784; 008857;
     02-FEB-2004, integrated into UniProtKB/Swiss-Prot.
     01-JUL-1997, sequence version 1.
DT
DT
     07-FEB-2006, entry version 37.
DE
     Treacle protein (Treacher Collins syndrome protein homolog).
GN
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os
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OC
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OC
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OC
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OX
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ŘΝ
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RP
     NUCLEOTIDE SEQUENCE [MRNA].
RC
     TISSUE=Liver;
RX
     MEDLINE=97445113; PubMed=9299440; DOI=10.1006/bbrc.1997.7229;
     Paznekas W.A., Zhang N., Gridley T., Jabs E.W.;
RT
      "Mouse TCOF1 is expressed widely, has motifs conserved in nucleolar
RT
     phosphoproteins, and maps to chromosome 18.";
     Biochem. Biophys. Res. Commun. 238:1-6(1997).
RN
     T21
RP
     NUCLEOTIDE SEQUENCE [MRNA], AND DEVELOPMENTAL STAGE.
     MEDLINE=97301769; PubMed=9158147; DOI=10.1093/hmg/6.5.727;
RA
     Dixon J., Hovanes K., Shiang R., Dixon M.J.;
RT
     "Sequence analysis, identification of evolutionary conserved motifs
RT
     and expression analysis of murine tcofl provide further evidence for a
RT
     potential function for the gene and its human homologue, TCOF1.";
RL
     Hum. Mol. Genet. 6:727-737(1997).
RN
     NUCLEOTIDE SEQUENCE [LARGE SCALE MRNA] OF 1-1314.
RP
RC
     STRAIN=C57BL/6; TISSUE=Brain;
RX
     MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA
     Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
     Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D., Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA
RA
     Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
     Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L., Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA
     Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C., Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA
     Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
     Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
     Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA
RA
     Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
     Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
     Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C., Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA
     Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
     Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
RA
RT
     "Generation and initial analysis of more than 15,000 full-length human
     and mouse cDNA sequences.";
RL
     Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN
     PHOSPHORYLATION SITE SER-1191.
RX
     PubMed=14729942; DOI=10.1074/mcp.D300003-MCP200;
RA
     Shu H., Chen S., Bi Q., Mumby M., Brekken D.L.;
     "Identification of phosphoproteins and their phosphorylation sites in
RT
     the WEHI-231 B lymphoma cell line.";
RL
     Mol. Cell. Proteomics 3:279-286(2004).
     -!- FUNCTION: May be involved in nucleolar-cytoplasmic transport. May
CC
CC
          play a fundamental role in early embryonic development,
CC
         particularly in development of the craniofacial complex.
     -!- SUBCELLULAR LOCATION: Nuclear; nucleolar (Potential).
CC
     -!- TISSUE SPECIFICITY: Ubiquitous in adult and embryonic tissues.
CC
     -!- DEVELOPMENTAL STAGE: Expression elevated at 11 dpc when the
         branchial arches and facial swellings are present.
CC
     -!- SIMILARITY: Contains 1 LisH domain.
CC
CC
     Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms
CC
     Distributed under the Creative Commons Attribution-NoDerivs License
CC
     EMBL; AF001794; AAB71347.1; -; mRNA.
DR
     EMBL; U81030; AAB60933.1; -; mRNA.
     EMBL; BC060105; AAH60105.1; -; mRNA.
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     GO; GO:0042790; P:transcription of nuclear rRNA large RNA pol. . .; IMP.
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     InterPro; IPR003993; treacle.
DR
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DR
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     SMART; SM00667; Lish; 1.
DR
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     Nuclear protein; Phosphorylation; Repeat; Transport.
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FT
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     COMPBIAS
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                         1314
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                          145
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Qу
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